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<!--StartFragment-->RESULT 7
US-08-385-229-2
; Sequence 2, Application US/08385229
; Patent No. 5605690
; GENERAL INFORMATION:
;   APPLICANT:  Jacobs, Cindy A.
;   APPLICANT:  Smith, Craig A.
;   TITLE OF INVENTION:  Method of Treating TNF-Dependent
;   TITLE OF INVENTION:  Inflammation Using Tumor Necrosis Factor Antagonists
;   NUMBER OF SEQUENCES:  5
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE:  Immunex Corporation
;     STREET:  51 University Street
;     CITY:  Seattle
;     STATE:  Washington
;     COUNTRY:  U.S.A.
;     ZIP:  98101
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE:  Floppy disk
;     COMPUTER:  IBM PC compatible
;     OPERATING SYSTEM:  PC-DOS/MS-DOS
;     SOFTWARE:  PatentIn Release #1.0, Version #1.25
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER:  US/08/385,229
;     FILING DATE:
;     CLASSIFICATION:  435
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:  US/07/946,236
;     FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;     NAME:  Wight, Christopher L.
;     REGISTRATION NUMBER:  31,680
;     REFERENCE/DOCKET NUMBER:  2503
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE:  (206) 587-0430
;     TELEFAX:  (206) 587-0606
;   INFORMATION FOR SEQ ID NO:  2:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  461 amino acids
;       TYPE:  amino acid
;       TOPOLOGY:  linear
;     MOLECULE TYPE:  protein
US-08-385-229-2

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Query Match          100.0%;  Score 1318;  DB 1;  Length 461;
Best Local Similarity 100.0%;  Pred. No. 5.8e-97;
Matches 235;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

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Qy      1  LPAQVAFTPYAPEPGSTCRLREYYDQTAQMCCSKCSPGQHAKVFCTKTSDTVCDSCEDST 60
        |||
Db      23  LPAQVAFTPYAPEPGSTCRLREYYDQTAQMCCSKCSPGQHAKVFCTKTSDTVCDSCEDST 82

Qy      61  YTQLWNWVPECLSCGSRSSDQVETQACTREQNRICTRPGWYCALSKEGCRLCAPLRK 120
        |||
Db      83  YTQLWNWVPECLSCGSRSSDQVETQACTREQNRICTRPGWYCALSKEGCRLCAPLRK 142

Qy     121  CRPGFGVARPGTETSDVVKPCAPGTFSNTTSSTDICRPHQICNVVAIPGNASMDAVCTS 180
        |||
Db     143  CRPGFGVARPGTETSDVVKPCAPGTFSNTTSSTDICRPHQICNVVAIPGNASMDAVCTS 202

Qy     181  TSPTRSMAPGAVHLPQPVSTRSQHTQPTPEPSTAPSTSFLLLPMGPSPPAEGSTGD 235

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Db          203  TSPTRSMAPGAVHLPQPVSTRSQHTQPTPEPSTAPSTSFLLLPMGPPPAEGSTGD 257

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